

- (ii) MOLECULE TYPE: peptide
- (v) FRAGMENT TYPE: N-terminal
- (vi) ORIGINAL SOURCE:
(A) ORGANISM: alkalophilic Bacillus sp.
(B) STRAIN: 38-2
- (vii) IMMEDIATE SOURCE:
(B) CLONE: beta-cyclodextrin
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Ala Pro Asp Thr Ser Val Ser Asn Lys Gln Asn Phe Ser Thr Asp Val
1 5 10 15

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REMARKS

The present Amendment is in response to Examiner's Supplemental Notice to Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures mailed August 9, 1993 and the original Notice to Comply mailed April 8, 1993.

Applicants submit this Amendment to complete their disclosure of the sequences employed in the present application pursuant to 37 C.F.R. § 1.821 et seq. Applicants also submit herewith a paper copy of the Sequence Listing and a floppy disk containing the Sequence Listing in computer readable form in compliance with 37 C.F.R. §§ 1.821 (c) and (e).

The Examiner indicates in the Notice that an error in the Sequence Listing exists with regard to sequence number 7 which is said to contain 709 bases although

only 708 bases are disclosed. Sequence 7 does contain 709 base pairs as is stated in the Sequence Listing. Rather, the error in the amended sequence listing submitted June 2, 1993 relates to the inadvertant omission of a single base "A" at position 143 of sequence number 7. This inadvertant error was only realized after the Examiner indicated that sequence number 7 disclosed 708 bases as opposed to the 709 bases it is said to contain. In response, Applicants have amended sequence number 7 so that bases 141-150 of sequence number 7 read "TGAAAAATTC" as opposed to "TGAAAATTC".

Entry of the present amendment to the Sequence Listing is proper since no new matter is being introduced by the present amendment to the Specification or by these submissions. Applicants originally disclosed that sequence number 7 contains 709 bases and inadvertently omitted a single base at position 143. This omission is evident upon inspection since sequence grouping 141-150 contains only 9 bases. Applicants' present amendment only adds the tenth base that was inadvertently omitted. Applicants maintain that the present amendment to sequence number 7 does not constitute new matter.

Entry of the present amendment to the sequence listing is also proper since the content of the amendment to the Specification and the paper and computer readable copies of the Sequence Listing are the same.

Respectfully submitted,
LIMBACH & LIMBACH

Dated: 9/29/93

By: Albert P. Halluin
Albert P. Halluin
Registration No. 25,227

Attorneys for Applicant